## SEQUENCE LISTING

- (1) GENERAL INFORMATION:
  - (i) APPLICANT: Higuchi, Russell H.
- (ii) TITLE OF INVENTION: Homogeneous Methods for Nucleic Acid

Amplification and Detection

- (iii) NUMBER OF SEQUENCES: 14
  - (iv) CORRESPONDENCE ADDRESS:
    - (A) ADDRESSEE: Cetus Corporation
    - (B) STREET: 1400 Fifty-Third Street
    - (C) CITY: Emeryville
    - (D) STATE: California
    - (E) COUNTRY: USA
    - (F) ZIP: 94608
    - (v) COMPUTER READABLE FORM:
      - (A) MEDIUM TYPE: Floppy disk
      - (B) COMPUTER: IBM PC compatible
      - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
      - (D) SOFTWARE: PatentIn Release #1.0, Version
- #1.25
  - (vi) CURRENT APPLICATION DATA:
    - (A) APPLICATION NUMBER:
    - (B) FILING DATE:
    - (C) CLASSIFICATION:
  - (viii) ATTORNEY/AGENT INFORMATION:
    - (A) NAME: Sias, Stacey R.
    - (B) REGISTRATION NUMBER: 32,630
    - (C) REFERENCE/DOCKET NUMBER: 2599
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      - (C) TELEX: 4992659

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PATENT Atty. Docket No. 2599

(2) INFORMATION FOR SEQ ID NO:1:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 26 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
GTGCTGCAGG TGTAAACTTG TACCAG	26
(2) INFORMATION FOR SEQ ID NO:2:	
<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 28 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
CACGGATCCG GTAGCAGCGG TAGAGTTG	28
(2) INFORMATION FOR SEQ ID NO:3:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 17 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
GTGTAAACTT GTACCAG	17



(2)	INFO	PRMATION FOR SEQ ID NO:4:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 16 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:4:	
GGTA	GCAG	CG GTAGAG	16
(2)	INFO	RMATION FOR SEQ ID NO:5:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 27 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
		MOLECULE TYPE: DNA (genomic)	
		SEQUENCE DESCRIPTION: SEQ ID NO:5:	
		TT CGTGTCCCCA CAGCACG	27
(2)	INFO	RMATION FOR SEQ ID NO:6:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 27 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
		MOLECULE TYPE: DNA (genomic)	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:6:	
CTCC	CCAAC	CC CCGTAGTTGT GTCTGCA	27



(2)	RMATION FOR SEQ ID NO:/:		
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 21 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:7:	
TCCA	CTTT	AT TCCAGGCCTG T	21
(2)	INFO	RMATION FOR SEQ ID NO:8:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
		MOLECULE TYPE: DNA (genomic)	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:8:	
TTGA	ATGG	AA TGGGAACGAA TGG	23
(2)	INFO	RMATION FOR SEQ ID NO:9:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 19 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA (genomic)	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:9:	
ΑΑΤΑ	GACC	AA TAGGCAGAG	1 a



(2) IN	2) INFORMATION FOR SEQ ID NO:10:		
(.	<ul> <li>i) SEQUENCE CHARACTERISTICS:         <ul> <li>(A) LENGTH: 14 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul> </li> </ul>	•	
(i.	i) MOLECULE TYPE: DNA (genomic)		
(x.	i) SEQUENCE DESCRIPTION: SEQ ID NO:10:		
CACCTG	ACTC CTGA	14	
(2) IN	FORMATION FOR SEQ ID NO:11:		
(:	i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 14 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
	i) MOLECULE TYPE: DNA (genomic) i) SEQUENCE DESCRIPTION: SEQ ID NO:11:		
CACCTG	ACTC CTGT	14	
(2) IN	FORMATION FOR SEQ ID NO:12:		
(:	i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 26 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
(i:	i) MOLECULE TYPE: DNA (genomic)		
(x:	i) SEQUENCE DESCRIPTION: SEQ ID NO:12:		
CCAGGC	CAGA TGAGAGAACC AAGGGG	26	



- (2) INFORMATION FOR SEQ ID NO:13:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 53 base pairs

    - (B) TYPE: nucleic acid(C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: DNA (genomic)
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

GCGGGCAGGG CGGCGGGGC GGGGCCGAAC CGGTCTACAT AGTCTCTAAA GGG 53

- (2) INFORMATION FOR SEQ ID NO:14:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 17 base pairs

    - (B) TYPE: nucleic acid (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: DNA (genomic)
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

GGTCCCTGTC TTATGTC

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